

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

#14
BD
6-25-02Applicant: *Braun, et al.*

Serial No. 09/687,483

Filed: October 13, 2000

For: METHODS FOR GENERATING
DATABASES AND DATABASES FOR
IDENTIFYING POLYMORPHIC
GENETIC MARKERS

Art Unit: 1645

Examiner: Unassigned

INFORMATION DISCLOSURE STATEMENT IN
ACCORDANCE WITH 37 C.F.R. §§ 1.97-1.98Assistant Commissioner for Patents
Washington, D.C. 20231

Dear Sir:

Because this Information Disclosure Statement is filed prior to receipt of a First Office Action on the Merits for the above-captioned application, a fee for filing this statement should not be due. If, however, it is determined that a fee is due, the Commissioner is authorized to charge the unpaid amount to Deposit Account No. 50-1213.

In accordance with the duty of disclosure imposed by 37 C.F.R. § 1.56 to inform the Patent Office of all references known by Applicant or Applicant's representative that may be material to the examination of the subject application, Applicant's representative hereby provides this Information Disclosure Statement that is prepared in accordance with 37 C.F.R. §§ 1.97-1.98. Forms PTO-1449 (12 pages) and cited references are provided herewith in connection with the above-captioned application.

The documents listed on the Forms PTO-1449 and supplied herewith are in the English language with exception of item CL. International Patent Application No. WO 99/50447 (item CL), which is in the German language, is provided with an English language abstract. Hence, in accordance with the requirements of 37 C.F.R. § 1.98, as amended effective March 16, 1992, no further explanation of the listed items is necessary.

While the applicant does not wish to convey that the following references are the

U.S.S.N. 09/687,483

Braun, et al.

IDS

most relevant among the references cited, the following references are called to the attention of the Office: Item nos. CL, CU, DX, FL, GL, GM, and GP.

Applicant also makes known to the Examiner the following co-pending U.S. and International applications that have one or more common inventors and/or one or more common owners:

<u>U.S.S.N.</u>	<u>Filing Date</u>
08/467,208	06/06/95
08/481,033	06/07/95
08/617,010	03/18/96
08/744,481	11/06/96
08/786,988	01/23/97
08/922,201	09/02/97
08/947,801	10/08/97
09/146,054	09/02/98
09/179,536	10/26/98
09/266,409	03/10/99
09/285,481	04/02/99
09/287,141	04/06/99
09/287,681	04/06/99
09/287,682	04/06/99
09/287,679	04/06/99
09/297,576	05/03/99
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09/297,611	05/03/99
09/337,388	06/21/99
09/355,705	02/04/98
09/364,774	07/30/99
09/371,150	08/09/99
09/397,766	09/15/99
09/429,683	10/28/99
09/431,613	11/02/99
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09/504,245	02/15/00
09/566,591	05/08/00
09/584,258	05/31/00
09/604,696	06/26/00
09/628,478	07/31/00
09/664,977	09/18/00
09/678,620	10/02/00
09/680,581	10/05/00
09/686,148	10/10/00
09/724,877	11/28/00
60/037,165	02/04/97
60/159,176	10/13/99
60/217,251	07/10/00
60/217,658	07/10/00
60/240,335	10/13/00

U.S.S.N. 09/687,483

Braun, et al.

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Int'l Appln.
PCT/US00/28413

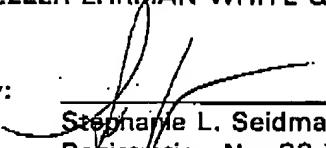
Filing Date
10/13/00

Although these documents and information are made known to the Patent and Trademark Office in compliance with Applicant's duty of disclosure, such disclosure is not to be construed as an admission by Applicant or Applicant's representative that any of the references, singly or in any combination thereof, is effective as prior art against the subject application. In accordance with 37 C.F.R. § 1.97(h), the filing of this Information Disclosure Statement shall not be construed to mean that a search has been made or that no other material information as defined in 37 C.F.R. § 1.56(b) exists.

Applicant respectfully requests that the Examiner review the foregoing references and they be made of record in the file history of the above-captioned application.

* * *

Respectfully submitted,
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Dated: February 15, 2001

Attorney Docket No. 24736-2033

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Sheet 1 of 12

FORM PTO-1449 (Modified)								ATTY. DOCKET NO. 24736-2033	SERIAL NO. 09/687,483
LIST OF PATENTS AND PUBLICATIONS FOR APPLICANT'S INFORMATION DISCLOSURE STATEMENT								APPLICANT Braun <i>et al.</i>	
								FILING DATE October 13, 2000	GROUP 1645

U.S. PATENT DOCUMENTS

EXAMINER INITIAL		DOCUMENT NUMBER							DATE	NAME	CLASS	SUB CLASS	FILING DATE
	AA	4	6	8	3	1	9	5	07/28/87	Mullis <i>et al.</i>	435	6	02/07/86
	AB	4	6	8	3	2	0	2	07/28/87	Mullis	435	91	10/25/85
	AC	4	8	2	6	3	6	0	05/02/89	Iwasawa <i>et al.</i>	406	51	02/25/87
	AD	4	8	5	1	0	1	8	07/25/89	Lazzari <i>et al.</i>	55	356	11/20/87
	AE	5	1	1	8	9	3	7	06/02/92	Hillenkamp <i>et al.</i>	250	282	08/21/90
	AF	5	4	3	6	1	5	0	07/25/95	Chandrasegaran	435	199	09/27/93
	AG	5	4	4	0	1	1	9	08/08/95	Labowsky	250	282	03/30/94
	AH	5	4	5	3	6	1	3	09/26/95	Gray <i>et al.</i>	250	281	10/21/94
	AI	5	4	9	8	5	4	5	03/12/96	Vestal	436	47	07/21/94
	AJ	5	5	0	3	9	8	0	04/02/96	Cantor	435	6	10/17/94
	AK	5	5	0	6	1	3	7	04/09/96	Mathur <i>et al.</i>	435	252.3	07/22/93
	AL	5	5	3	6	6	4	9	07/16/96	Fraiser <i>et al.</i>	435	91.2	07/29/94
	AM	5	5	4	7	8	3	5	08/20/96	Koster <i>et al.</i>	435	6	01/06/94
	AN	5	6	0	4	0	9	8	02/18/97	Mead <i>et al.</i>	435	6	12/22/94
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	AS	5	7	0	0	6	7	2	12/23/97	Mathur <i>et al.</i>	435	183	07/23/92
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	AW	5	8	4	3	6	6	9	12/01/98	Kaiser <i>et al.</i>	435	6	11/29/96
	AX	5	8	5	1	7	6	5	12/22/98	Koster	435	6	05/30/95

EXAMINER

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Sheet 2 of 12

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								FILING DATE October 13, 2000	GROUP 1645

U.S. PATENT DOCUMENTS

EXAMINER INITIAL		DOCUMENT NUMBER							DATE	NAME	CLASS	SUB CLASS	FILING DATE
	AY	5	8	5	8	7	0	5	01/12/99	Wei <i>et al.</i>	435	69.1	06/05/95
	AZ	5	8	7	1	9	1	1	02/16/99	Dahlberg <i>et al.</i>	435	6	02/09/95
	BA	5	8	7	2	0	0	3	02/16/99	Köster	435	283.1	05/30/95
	BB	5	8	7	4	2	8	3	02/23/99	Harrington <i>et al.</i>	435	252.3	05/30/95
	BC	5	8	8	5	8	4	1	03/23/99	Higgs, Jr. <i>et al.</i>	436	89	09/11/96
	BD	5	8	8	8	7	9	5	05/30/99	Hamilton	435	200	09/09/97
	BE	5	9	0	0	4	8	1	05/04/99	Lough <i>et al.</i>	536	55.3	11/06/96
	BF	5	9	2	8	9	0	6	07/27/99	Koster <i>et al.</i>	435	91.2	05/09/96
	BG	5	9	5	2	1	7	6	09/14/99	McCarthy <i>et al.</i>	435	6	12/21/95
	BH	5	9	7	6	8	0	6	11/02/99	Mahajan <i>et al.</i>	435	6	06/27/98
	BI	6	0	2	2	6	8	8	02/08/00	Jurinka <i>et al.</i>	435	6	06/13/96
	BJ	6	0	2	4	9	2	5	02/15/00	Little <i>et al.</i>	422	100	01/23/97
	BK	6	0	4	3	0	3	1	03/28/00	Koster <i>et al.</i>	435	6	03/18/96
	BL	6	0	5	4	2	7	6	04/25/00	Macevick	435	6	02/23/98
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	BN	6	0	9	0	6	0	6	07/18/00	Kaiser <i>et al.</i>	435	199	12/02/96
	BO	6	0	9	9	5	5	3	08/08/00	Hart <i>et al.</i>	606	232	05/21/98
	BP	6	1	3	3	4	3	6	10/17/00	Koster <i>et al.</i>	536	24.3	09/19/97
	BQ	6	1	4	0	0	5	3	10/31/00	Koster <i>et al.</i>	435	6	09/25/98
	BR	6	1	4	6	8	5	4	11/14/00	Koster <i>et al.</i>	435	91.1	08/31/95

FOREIGN PATENT DOCUMENTS

		DOCUMENT NUMBER							DATE	COUNTRY	CLASS	SUB CLASS	Translation Yes	No
	BS	0	0	5	6	4	4	6	09/28/00	PCT				

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Sheet 3 of 12

FORM PTO-1449 (Modified)								ATTY. DOCKET NO. 24736-2033	SERIAL NO. 09/687,483
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FOREIGN PATENT DOCUMENTS

		DOCUMENT NUMBER							DATE	COUNTRY	CLASS	SUB CLASS	Translation Yes No
	BT	0	0	6	0	3	6	1	11/12/00	PCT			
	BU	0	5	9	6	2	0	5	05/11/94	EP			
	BV	2	7	4	9	6	8	2	12/12/97	FR			
	BW	9	3	1	5	4	0	7	08/05/93	PCT			
	BX	9	4	1	6	1	0	1	07/21/94	PCT			
	BY	9	4	2	1	8	2	2	09/29/94	PCT			
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	CA	9	7	0	3	2	1	0	01/30/97	PCT			
	CB	9	7	0	8	3	0	6	03/06/97	PCT			
	CC	9	7	3	7	0	4	1	10/09/97	PCT			
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	CG	9	8	2	0	0	2	0	05/14/98	PCT			
	CH	9	8	2	0	1	6	6	05/14/98	PCT			
	CI	9	8	3	3	8	0	8	08/06/98	PCT			
	CJ	9	9	1	2	0	4	0	03/11/99	PCT			
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	CL	9	9	5	0	4	4	7	10/07/99	PCT			
	CM	9	9	6	4	5	0	1	10/28/99	PCT			
	CN	9	9	5	7	3	1	8	11/11/99	PCT			

OTHER ART (Including Author, Title, Date, Pertinent Pages, Etc.)

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Sh et 4 of 12

LIST OF PATENTS AND PUBLICATIONS FOR APPLICANT'S INFORMATION DISCLOSURE STATEMENT	ATTY. DOCKET NO. 24736-2033	SERIAL NO. 09/687,483
	APPLICANT Braun <i>et al.</i>	
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	CO	Arrand <i>et al.</i> , Different Substrate Specificities of the Two DNA Ligases of Mammalian Cells, <i>J. Biol. Chem.</i> 261(20):9079-82 (1986).
	CP	Badger <i>et al.</i> , New features and enhancements in the X-PLOR computer program, <i>Proteins: Structure, Function, and Genetics</i> 35(1):25-33 (1999)
	CQ	Beck <i>et al.</i> , Chemiluminescent detection of DNA: application for DNA sequencing and hybridization, <i>Nucl. Acids Res.</i> 17(13):5115-23 (1989).
	CR	Bertina <i>et al.</i> , Mutation in blood coagulation factor V associated with resistance to activated protein C, <i>Nature</i> 369:647 (1994).
	CS	Bessho <i>et al.</i> , Nucleotide excision repair 3' endonuclease XPG stimulates the activity of base excision repair enzyme thymine glycol DNA glycosylase, <i>Nucl. Acids Res.</i> 27(4):79-83 (1999).
	CT	Bjelland, S. and E. Seeberg, Purification and characterization of 3-methyladenine DNA glycosylase I from <i>Escherichia coli</i> , <i>Nucl. Acids Res.</i> 15(7):2787-2800 (1987).
	CU	Bleczinski, C. and C. Richert, Monitoring the Hybridization of the Components of Oligonucleotide Mixtures to Immobilized DNA via Matrix-assisted Laser Desorption/Ionization Time-of-flight Mass Spectrometry, <i>Rapid Communications in Mass Spectrometry</i> 12:1737-43 (1998).
	CV	Braun <i>et al.</i> , Detecting <i>CFTR</i> gene mutations by using primer oligo base extension and mass spectrometry, <i>Clinical Chemistry</i> 43(7):1151-8 (1997).
	CW	Braun <i>et al.</i> , Improved Analysis of Microsatellites Using Mass Spectrometry, <i>Genomics</i> 46:18-23 (1997).
	CX	Bregman <i>et al.</i> , Molecular Characterization of Bovine Brain P75, a High Affinity Binding Protein for the Regulatory Subunit of cAMP-dependent Protein Kinase II β , <i>J. Biol. Chem.</i> 266(11):7207-13 (1991).
	CY	Buetow <i>et al.</i> , High-throughput development and characterization of a genome-wide collection of gene-based single nucleotide polymorphism markers by chip-based matrix-assisted laser desorption/ionization time-of-flight mass spectrometry, <i>Proc. Natl. Acad. Sci. USA</i> 98(2):581-4 (2001).
	CZ	Burton <i>et al.</i> , Type II regulatory subunits are not required for the anchoring-dependent modulation of Ca^{2+} channel activity by cAMP-dependent protein kinase, <i>Proc. Natl. Acad. Sci. USA</i> 94:11067-72 (1997).
	DA	Carr <i>et al.</i> , Association of the Type II cAMP-dependent Protein Kinase with a Human Thyroid RII-anchoring Protein, <i>J. Biol. Chem.</i> 267(19):13376-82 (1992).

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	DB	Carr <i>et al.</i> , Interaction of the Regulatory Subunit (RII) of cAMP-dependent Protein Kinase with RII-anchoring Protein Occurs through an Amphipathic Helix Binding Motif, <u>J. Biol. Chem.</u> 266(22):14188-92 (1991).
	DC	Chiu <i>et al.</i> , Mass Spectrometry of Nucleic Acids, <u>Clin. Chem.</u> 45:1578 (1999).
	DD	Chiu <i>et al.</i> , Mass Spectrometry of single-stranded restriction fragments captured by an undigested complementary sequence, <u>Nucl. Acids. Res.</u> 28(8):e31 (2000).
	DE	Clegg <i>et al.</i> , Genetic characterization of a brain-specific form of the type I regulatory subunit of cAMP-dependent protein kinase, <u>Proc. Natl. Acad. Sci. USA</u> 85:3703-7 (1988).
	DF	Coghlan <i>et al.</i> , Association of Protein Kinase A and Protein Phosphatase 2B with a Common Anchoring Protein, <u>Science</u> 267:108-111 (1995).
	DG	Colledge, M and J.D. Scott., AKAPs: from structure to function, <u>Trends in Cell Biology</u> 9:216-21 (1999).
	DH	Corder <i>et al.</i> , Gene Dose of Apolipoprotein E Type 4 Allele and the Risk of Alzheimer's Disease in Late Onset Families, <u>Science</u> 261:921-3 (1993).
	DI	Database WPI, Derwent publication # 011635345 citing International Patent Application WO 9747974 of the parent French Patent Application FR 2,749,662.
	DJ	Eftedal <i>et al.</i> , Consensus sequences for good and poor removal of uracil from double stranded DNA by uracil-DNA glycosylase, <u>Nucl. Acids Res.</u> 21(9):2095-101 (1993).
	DK	Faux, M.C. and J.D. Scott., More on target with protein phosphorylation: conferring specificity by location, <u>Trends Biochem</u> 21:312-5 (1996).
	DL	Fu <i>et al.</i> , Efficient preparation of short DNA sequence ladders potentially suitable for MALDI-TOF DNA sequencing, <u>Genetic Analysis: Biomolecular Engineering</u> 12:137-42 (1996).
	DM	Fu <i>et al.</i> , Sequencing Exons 5 to 8 of the p53 Gene by MALDI-TOF Mass Spectrometry, <u>Nature Biotechnol.</u> 16:381-4 (1998).
	DN	Fu <i>et al.</i> , A DNA sequencing strategy that requires only five bases of known terminal sequence for priming, <u>Proc. Natl. Acad. Sci. USA</u> 92:10162-66 (1995).
	DO	Fu <i>et al.</i> , Sequencing double-stranded DNA by strand displacement, <u>Nucl. Acids Res.</u> 25(3):677-9 (1997).
	DP	Gabbita <i>et al.</i> , Decrease in Peptide Methionine Sulfoxide Reductase in Alzheimer's Disease Brain, <u>J. Neurochemistry</u> 73(4):1660-6 (1999).

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Sheet 6 of 12

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LIST OF PATENTS AND PUBLICATIONS FOR APPLICANT'S INFORMATION DISCLOSURE STATEMENT		APPLICANT Braun <i>et al.</i>	
		FILING DATE October 13, 2000	GROUP 1645

	DQ	Giantz <i>et al.</i> , Characterization of Distinct Tethering and Intracellular Targeting Domains in AKAP75, a Protein That Links cAMP-dependent Protein Kinase II β to the Cytoskeleton, <u>J. Biol. Chem.</u> 268(17):12796-804 (1993).
	DR	Goldmacher <i>et al.</i> , Photoactivation of toxin conjugates, <u>Bioconj. Chem.</u> 3:104-107 (1992)
	DS	Guatelli <i>et al.</i> , Isothermal, <i>in vitro</i> amplification of nucleic acids by a multienzyme reaction modeled after retroviral replication, <u>Proc. Natl. Acad. Sci. USA</u> 87:1874-8 (1990).
	DT	Hausken <i>et al.</i> , Mutational Analysis of the A-Kinase Anchoring Protein (AKAP)-binding Site on RII, <u>J. Biol. Chem.</u> 271(46):29016-22 (1996).
	DU	Hazum <i>et al.</i> , A photocleavable protecting group for the thiol function of cysteine, in <u>Pept., Proc. Eur. Pept. Symp., 16th</u> Brunfeldt, K (ed), pp. 105-110- (1981)
	DV	Higgins <i>et al.</i> , Competitive Oligonucleotide Single-Base Extension Combined with Mass Spectrometric Detection for Mutation Screening, <u>BioTechniques</u> 23(4):710-4 (1997).
	DW	Higgins <i>et al.</i> , DNA-Joining Enzymes: A Review, <u>Methods in Enzymology</u> 68:50-71 (1979).
	DX	Higley <i>et al.</i> , Processivity of uracil DNA glycosylase, <u>Mutation Research, DNA Repair</u> 294:109-116 (1993).
	DY	Hinton <i>et al.</i> , The application of robotics to fluorometric and isotopic analyses of uranium, <u>Laboratory Automation & Information Management</u> , NL, Elsevier Science publishers BV., Amsterdam, Vol. 21 no. 2/03, pp. 223-227, December 1, 1993.
	DZ	Huang <i>et al.</i> , D-AKAP2, a novel protein kinase A anchoring protein with a putative RGS domain, <u>Proc. Natl. Acad. Sci. USA</u> 94:11184-9 (1997).
	EA	Hubbard, M.J. and P. Cohen., On target with a new mechanism for the regulation of protein phosphorylation, <u>Trends Biochem. Sci.</u> 18:172-77 (1993).
	EB	Instrumentation; "Nano-Plotter" from GeSiM, Germany, located at http://www.gesim.de/np-intro.htm
	EC	Instrumentation; "Model CRS A 255" robot "Digital Servo Gripper" "Plate Cube" system. "lid parking station" "shaker" Robocon Labor-und Industrieroboter Ges.m.b.H of Austria ("Robocon")

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Sheet 7 of 12

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	ED	Instrumentation: "MJ Microseal" plate sealer; Thermal Cycler Accessories: Sealing Options, Sealing Products, MJ Research, located at http://www.mjresearch.com/html/consumables/ealing/sealing_products.html
	EE	Instrumentation: "Genesis 200/8" (200 cm with including an 8-tip arm) liquid handling systems; Tecan AG of Switzerland ("Tecan"), TECAN Products for Diagnostics and Life Science, located at http://www.tecan.ch/index.htm
	EF	Instrumentation; Bar code systems, including one and two dimensional bar codes, readable and readable/writable codes and systems; Datalogic S.p.A. of Italy ("Datalogic") located at http://www.datalogic.com
	EG	Instrumentation; DYNABEADS, streptavidin-coated magnetic beads; from Dynal, Inc. Great Neck, NY and Oslo Norway
	EH	Instrumentation; "Multimek 96" automated pipettor; Beckman Coulter, Inc. located at http://www.coulter.com , 09/08/99
	EI	International Search Report for International Application No. PCT/US00/08111, Date of Mailing November 13, 2000.
	EJ	Jahnsen <i>et al.</i> , Molecular Cloning, cDNA Structure, and Regulation of the Regulatory Subunit of Type II cAMP-dependent Protein Kinase from Rat Ovarian Granulosa Cells, <u>J. Biol. Chem.</u> 261(26):12352-61 (1986).
	EK	Jurinke <i>et al.</i> , Recovery of Nucleic Acids from Immobilized Biotin-Streptavidin Complexes Using Ammonium Hydroxide and Applications in MALDI-TOF Mass Spectrometry, <u>Anal. Chem.</u> 69:904-10 (1997).
	EL	Jurinke <i>et al.</i> , Analysis of Ligase Chain Reaction products via Matrix-Assisted Laser Desorption/Ionization Time-of-Flight-Mass Spectrometry, <u>Anal. Biochem.</u> 237:174-81 (1996).
	EM	Jurinke <i>et al.</i> , Detection of hepatitis B virus DNA in serum samples via nested PCR and MALDI-TOF mass spectrometry, <u>Genetic Analysis: Biomolecular Engineering</u> 13:67-71 (1996).
	EN	Jurinke <i>et al.</i> , Application of nested PCR and mass spectrometry for DNA-based virus detection: HBV-DNA detected in the majority of isolated anti-HBc positive sera, <u>Genetic Analysis: Biomolecular Engineering</u> 14:97-102 (1998).
	EO	Kario <i>et al.</i> , Genetic Determinants of Plasma Factor VII Activity in the Japanese, <u>Thromb. Haemost.</u> 73:617-22 (1995).
	EP	Klauck <i>et al.</i> , Coordination of Three Signaling Enzymes by AKAP79, a Mammalian Scaffold Protein, <u>Science</u> 271:1589-92 (1996).

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LIST OF PATENTS AND PUBLICATIONS FOR APPLICANT'S INFORMATION DISCLOSURE STATEMENT		APPLICANT Braun <i>et al.</i>	
		FILING DATE October 13, 2000	GROUP 1645

	EQ	Koster et al., Oligonucleotide synthesis and multiples DNA sequencing using chemiluminescent detection, <u>Nucl. Acids Res. Symposium Series No. 24</u> (1991) 318-21.	
	ER	Koster et al., A strategy for rapid and efficient DNA sequencing by mass spectrometry, <u>Nature Biotechnology</u> 14:1123-8 (1996).	
	ES	Kwoh et al., Transcription-based amplification system and detection of amplified human immunodeficiency virus type 1 with a bead-based sandwich hybridization format, <u>Proc. Natl. Acad. Sci. USA</u> 86:1173-7 (1989).	
	ET	Laken et al., Familial colorectal cancer in Ashkenazim due to a hypermutable tract in APC, <u>Nature Genetics</u> 17:79-83 (1995).	
	EU	Lam et al., Genetic influence of the R/Q353 genotype on factor VII activity is overwhelmed by environmental factors in Chinese patients with Type II (non-insulin-dependent) diabetes mellitus, <u>Diabetologia</u> 41:760-66 (1998).	
	EV	<u>Lasko et al., Eukaryotic DNA Ligases, Mutation Research</u> 236:277-87 (1990).	
	EW	Lee et al., Isolation of a cDNA clone for the type I regulatory subunit of bovine cAMP-dependent protein kinase, <u>Proc. Natl. Acad. Sci. USA</u> 80:3608-12 (1983).	
	EX	Lehman, I.R., DNA Ligase: Structure, Mechanism, and Function, <u>Science</u> 186:790-7 (1974).	
	EY	Li et al., DNA ligase 1 is associated with the 21 S complex of enzymes for DNA synthesis in HeLa cells, <u>Nucl. Acids Res.</u> 22(4):632-8 (1994).	
	EZ	Li et al., High-Resolution MALDI Fourier Transform Mass Spectrometry of Oligonucleotides, <u>Anal. Chem.</u> 68(13):2090-6 (1996).	
	FA	Lindahl, T. and D.E. Barnes., Mammalian DNA Ligases, <u>Annu. Rev. Biochem.</u> 61:251-81 (1992).	
	FB	Little et al., Detection of <i>RET</i> proto-oncogene codon 634 mutations using mass spectrometry, <u>J. Mol. Med.</u> 75:745-50 (1997).	
	FC	Little et al., Identification of apolipoprotein E polymorphisms using temperature cycled primer oligo base extension and mass spectrometry, <u>Eur J Clin Chem Clin Biochem</u> 35(7):545-8 (1997)	
	FD	Little et al., MALDI on a chip: analysis of arrays of low-femtomole to subfemtomole quantities of synthetic oligonucleotides and DNA diagnostic products dispensed by a piezoelectric pipet, <u>Anal. Chem.</u> 69:4540-4546 (1997)	
	FE	Little et al., Mass Spectrometry from miniaturized arrays for full comparative DNA analysis, <u>Nature Medicine</u> 3(12):1413-6 (1997).	

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Sheet 9 of 12

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	FF	Lizardi <i>et al.</i> , Exponential Amplification of Recombinant-RNA Hybridization Probes, <u>Bio/Technology</u> 6:1197-1202 (1988).
	FG	Miki, K. and E.M. Eddy, Single Amino Acids Determine Specificity of Binding Protein Kinase A Regulatory Subunits by Protein Kinase A Anchoring Proteins, <u>J. Biol. Chem.</u> 274(41):29057-62 (1999).
	FH	Miki, K. and E.M. Eddy, Identification of Tethering Domains for Protein Kinase A Type I α Regulatory Subunits on Sperm Fibrous Sheath Protein FSC1, <u>J. Biol. Chem.</u> 273(51):34384-90 (1996).
	FI	Mochly-Rosen, D., Localization of Protein Kinases by Anchoring Proteins: A Theme in Signal Transduction, <u>Science</u> 268:247-51 (1995).
	FJ	Moskovitz <i>et al.</i> , Overexpression of peptide-methionine sulfoxide reductase in <u>Saccharomyces cerevisiae</u> and human T cells provides them with high resistance to oxidative stress, <u>Proc. Natl. Acad. Sci. USA</u> 95:14071-5 (1998).
	FK	Nilges <i>et al.</i> , Automated NOESY interpretation with ambiguous distance restraints: the refined NMR solution structure of the pleckstrin homology domain from β -spectrin, <u>J. Mol. Biol.</u> 269:408-422 (1997).
	FL	<u>Nucleases</u> 2nd ed. Linn, S.M. <i>et al.</i> (eds.) Cold Spring Harbor Laboratory Press (1993).
	FM	Podhajska, A.J. and W. Szybalski, Conversion of the <i>FokI</i> endonuclease to a universal restriction enzyme: cleavage of phage M13mp7 DNA at predetermined sites, <u>Gene</u> 40:175-82 (1985).
	FN	Reymer <i>et al.</i> , A lipoprotein lipase mutation (Asn291Ser) is associated with reduced HDL cholesterol levels in premature atherosclerosis, <u>Nature Genetics</u> 10:28-34 (1995).
	FO	Ruppert <i>et al.</i> , A Filtration Method for Plasmid Isolation Using Microtiter Filter Plates, <u>Anal. Biochem.</u> 230:130-4 (1995).
	FP	Samson <i>et al.</i> , Resistance to HIV-1 infection in caucasian individuals bearing mutant alleles of the CCR-5 chemokine receptor gene, <u>Nature</u> 382:722-5 (1996).
	FQ	Saparbaev <i>et al.</i> , <i>Escherichia coli</i> , <i>Saccharomyces cerevisiae</i> , rat and human 3-methyladenine DNA glycosylases repair 1, N ⁶ -ethenoadenine when present in DNA, <u>Nucl. Acids Res.</u> 23(18):3750-55 (1995).
	FR	Schachter <i>et al.</i> , Genetic associations with human longevity at the APOE and ACE loci, <u>Nature Genetics</u> 6:29-32 (1994).
	FS	Scott <i>et al.</i> , Type II Regulatory Subunit Dimerization Determines the Subcellular Localization of the cAMP-dependent Protein Kinase, <u>J. Biol. Chem.</u> 265(35):21561-66 (1990).

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FT	Scott, J., Cyclic Nucleotide-Dependent Protein Kinases, <u>Pharmac. Ther</u> 50:123-45 (1991).
FU	Senko <i>et al.</i> , Automated Assignment of Charge States from Resolved Isotopic Peaks for Multiply Charged Ions, <u>J. Am. Soc. Mass Spectrom</u> 6:52-56 (1995).
FV	Senter <i>et al.</i> , Novel photocleavable protein crosslinking reagents and their use in the preparation of antibody-toxin conjugates, <u>Photachem. Photobiol.</u> 42:231-237 (1985)
FW	Sequenom Advances the Industrial Genomics Revolution with the Launch of Its DNA MassArray™ Automated Process Line, Press Release: Sept. 28, 1998, http://www.sequenom.com/pressrelease.htm .
FX	Sequenom Completes Design of More Than 400,000 SNP Assays; Mass EXTEND™ Assay Portfolio Covers Majority of SNPs in the Public Domain, Press Release; Oct. 10, 2000, http://www.sequenom.com/ir/ir_prs.asp
FZ	Sequenom and Gemini Identify Genes Linked to Cardiovascular Disease, Press Release: Nov. 28, 2000, http://www.sequenom.com/ir/ir_prs.asp
GA	Sequenom Announces Publication of Results From Large-Scale SNP Study With the National Cancer Institute, Press Release: Jan. 16, 2001, http://www.sequenom.com/ir/ir_prs.asp
GB	Sequenom: Technologies and Tools, located at http://www.sequenom.com/tech/tools.html , dated 08/29/99
GC	Siegert <i>et al.</i> , Matrix-Assisted Laser desorption/Ionization Time-of-Flight Mass Spectrometry for the detection of Polymerase Chain Reaction Containing 7-Deazapurine Moieties, <u>Anal. Biochem.</u> 243:55-65 (1996).
GD	Smith, L.M., Sequence from spectrometry: A realistic prospect, <u>Nature Biotechnology</u> 14:1084-5 (1996).
GE	Sugisaki, H. and S. Kanazawa, New restriction endonucleases from <i>Flavobacterium okeanokoites</i> (FokI) and <i>Micrococcus luteus</i> (MluI), <u>Gene</u> 16:73-8 (1981).
GF	Szybalski <i>et al.</i> , Class-IIIS restriction enzymes - a review, <u>Gene</u> 100:13-26 (1991).
GG	Takio <i>et al.</i> , Primary structure of the regulatory subunit of type II cAMP dependent protein kinase from bovine cardiac muscle, <u>Proc. Natl. Acad. Sci. USA</u> 79:2544-8 (1982).
GH	Tammnen <i>et al.</i> , Proteolytic cleavage of glucagon-like peptide-1 by pancreatic β cells and by fetal calf serum analyzed by mass spectrometry, <u>J. Chromatogr. A</u> 852:285-295 (1999).
GI	Tang <i>et al.</i> , Matrix-assisted laser desorption/ionization mass spectrometry of immobilized duplex DNA probes, <u>Nucl. Acids Res.</u> 23(16):3126-31 (1995).

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GJ	Tang <i>et al.</i> , Chip-based genotyping by mass spectrometry, <u>Proc. Natl. Acad. Sci. USA</u> 96:10016-20 (1999).
GK	Thompson, Fitting robots with white coats, <u>Laboratory Automation and Information Management</u> 31:173-193 (1996).
GL	Uracil-DNA Glycosylase, product description: Roche Molecular Biochemicals Catalog Version 3, Nov. 1999 http://biochem.roche.com/pack-insert/1269062a.pdf (12/21/00).
GM	Uracil-DNA Glycosylase (UDG), product description. New England Biolabs. http://circuit.neb.com/neb/products/mod_enzymes/280.html (12/21/00).
GN	van den Boom <i>et al.</i> , Forward and Reverse DNA Sequencing in a Single Reaction, <u>Anal. Biochem.</u> 256:127-9 (1998).
GO	van den Boom <i>et al.</i> , Combined amplification and sequencing in a single reaction using two DNA polymerase with differential incorporation rates for dideoxynucleotides, <u>J. Biochem. Biophys. Methods</u> 35:69-79 (1997).
GP	Vaughan <i>et al.</i> , Glycosylase mediated polymorphism detection (GMPD) – a novel process for genetic analysis, <u>Genetic Analysis: Biomolecular Engineering</u> 14:169-75 (1999).
GQ	Waga <i>et al.</i> , Reconstitution of Complete SV40 DNA Replication with Purified Replication Factors, <u>J. Biol. Chem.</u> 269(14):10923-34 (1994).
GR	Wang <i>et al.</i> , Allene γ_9 and γ_{10} : low-temperature measurements of line intensity, <u>J. Mol. Spectrosc.</u> 194(20):256-268 (1999).
GS	Weiler <i>et al.</i> , Hybridisation based DNA screening on peptide nucleic acid (PNA) oligomer arrays, <u>Nucleic Acids Res.</u> 25:2792-2799 (1997).
GT	Wilson <i>et al.</i> , Restriction and Modification Systems, <u>Annu. Rev. Genet.</u> 25:585-627 (1991).
GU	Yen <i>et al.</i> , Synthesis of water-soluble copolymers containing photocleavable bonds, <u>Makromol. Chem.</u> 190:69-82 (1989).

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